



ENTERED

RAW SEQUENCE LISTING

DATE: 03/20/2002 P.5

PATENT APPLICATION: US/10/082,830

3 <110> APPLICANT: Sun, Yongming

TIME: 11:47:04

Input Set : A:\Dex-249.app

Output Set: N:\CRF3\03202002\J082830.raw

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Recipon, Herve
              Salceda, Susana
              Liu, Chenghua
              Turner, Leah
      9 <120> TITLE OF INVENTION: Compositions and Methods Relating to Breast Specific
              Genes and Proteins
     12 <130> FILE REFERENCE: DEX-0249
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/082,830
C--> 15 <141> CURRENT FILING DATE: 2001-10-29
     17 <150> PRIOR APPLICATION NUMBER: 60/243,802
     18 <151> PRIOR FILING DATE: 2000-10-27
     20 <160> NUMBER OF SEQ ID NOS: 282
     22 <170> SOFTWARE: PatentIn Ver. 2.1
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     25 <211> LENGTH: 207
     26 <212> TYPE: DNA
     27 <213> ORGANISM: Homo sapiens
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     31 catttateca etggtaetat ttaqttqtat ttattaqaea qetteetgee etetecaaaa 120
     32 agcttactga gctagtaact atttacaggg ttagccaaag aacacaaaaa agtgatctct 180
     33 attagactgt aagaatatgg tttccct
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     42 <221> NAME/KEY: unsure
     43 <222> LOCATION: (492)
     44 <223> OTHER INFORMATION: a, c, g or t
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     49 accagtgttt coctagtact taacatgggt ttattacatt tttttgacaa aaattcaaaa 180
     50 ttacatatat tttgttette attageaagt cacacatttt aaaatggeae acteeettee 240
     51 tettegtgtt gttatttgtt tattttaagg aetgtttetg ggtagataag etetgggtta 300
     52 ttttaaaata cattttacaa tggaaatggc ctggacttga actgaaaagg aaacattatc 360
     53 tgtgttattt cagacacatc agtgatcagt ttagaagata ggatgatttc actaagctta 420
     54 taattcatet taaageteae etaaataaaa gtaagtgaet aaaatgatet tttteetteea 480
W--> 55 ggagaggtag gnttaattaa ttg
     58 <210> SEQ ID NO: 3
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59 <211> LENGTH: 603

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60 <212> TYPE: DNA
     61 <213> ORGANISM: Homo sapiens
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     66 accagtgttt ccctagtact taacatgggt ttattacatt tttttgacaa aaattcaaaa 180
     67 ttacatatat tttgttcttc attagcaagt cacacatttt aaaatggcac actcccttcc 240
     68 tettegtgtt gttatttgtt tattttaagg aetgtttetg ggtagataag etetgggtta 300
     69 ttttaaaata cattttacaa tggaaatggc ctggacttga actgaaaagg aaacattatc 360
     70 tgtgttattt cagacacatc agtgatcagt ttagaagata ggatgatttc actaagctta 420
     71 taattcatct taaagctcac ctaaataaaa qtaaqtqact aaaatqatct ttttcttcca 480
     72 ggagaggtag gattaattaa tggtataatg tgtggaatat ttcaggctta tctgattctt 540
     73 ccatcttaaa tctttgagag ttttaaacac attatgtgtc cattactgtt tatatcacat 600
     74 aga
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     83 <221> NAME/KEY: unsure
     84 <222> LOCATION: (133)
     85 <223> OTHER INFORMATION: a, c, g or t
     87 <400> SEQUENCE: 4
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     89 tgtcattttc tgggcacaga aagtgcgatg gaagggaatg agaaggggaa aaaggaagga 120
W--> 90 tgacaggacg ganggaggga aagaaggaag aggaaaaaag aaaggacagg agaaagggag 180
     91 gaaggettet gecaaaaaat taaaateaaa tttttgacat tetttttgtt tgeetttttt 240
     92 gaaacaaaat gacacttgcc agacaccagc ttcctggccc atgtcctggt ccttggtatc 300
     93 cagatgacag cagtgtgatc ctgctgtgag ttccttccgt gccttctgat ctgagttcct 360
     94 gaaagcaqaq agccactcaq gaactqctqt ctctcaqqcc aqctqqctqq tqatqqqctt 420
     95 ttgaaqactc tqqctctctc tcctqctqqa aqaqctcccc aqqqqccacc aqqaqccaqq 480
     96 tgaccgctct cagcctctgt gagctactgg agatcaccag accttcccac atcc
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     102 <213> ORGANISM: Homo sapiens
     104 <220> FEATURE:
     105 <221> NAME/KEY: unsure
     106 <222> LOCATION: (329)
     107 <223> OTHER INFORMATION: a, c, g or t
     109 <400> SEQUENCE: 5
    110 atgactetta acqaqeatge tgeetteaaa catetgttta acqaaqeaca tettgeaceq 60
    111 cccttaatcc atttaaccct gagtggacac agcacatgtt tcagagagca cagggttggg 120
    112 gcactgtgcc agacactgga gataataaag aaaaacagca ctgggcctat aattgggagt 180
    113 ctagatatac gatatggaaa tgcagctgac aatgcaagga gcaagaggac tcgcacagtg 240
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114 gtgcatggca gcttgctgtc attttctggg cacagaaagt gcgatggaag ggaatgagaa 300 W--> 115 ggggaaaag gaaggatgac aggacggang gagggaaaga aggaaggaag 360 116 gacaggagaa agggaggaag gcttctgcca aaaaattaaa atcaaatttt tgacattctt 420 117 tttgtttgcc ttttttgaaa caaaatgaca cttgccagac accagcttcc tggcccatgt 480

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118 cctggtcctt ggtatccaga tgacagcagt gtgatcctgc tgtgagttcc ttccgtgcct 540 119 tetgatetga gtteetgaaa geagagagee aeteaggaae tgetgtetet eaggeeaget 600 120 ggctggtgat gggcttttga agactctggc tctctctct gctggaagag ctccccaggg 660 121 gccaccagga gccaggtgac cgctctcagc ctctgtgagc tactggagat caccagacct 720 122 teceacatee egggeaggtg ceagggeett taaggagget ttetgetetg eagggatgtt 780 123 ctqtqqqctc caqtattctq qcqaqcatca qcttattctc qqcttaqtct tcttqctcta 840 124 tageteetge etetgttttg ettetttttg gtgatgeetg ttteacetta tgaagaggge 900 125 ctgtaaaatc caagctctgc acaaaccc 128 <210> SEO ID NO: 6 129 <211> LENGTH: 368 130 <212> TYPE: DNA 131 <213> ORGANISM: Homo sapiens 133 <220> FEATURE: 134 <221> NAME/KEY: unsure 135 <222> LOCATION: (305) 136 <223> OTHER INFORMATION: a, c, g or t 138 <400> SEQUENCE: 6 139 tattaagtga ctttaatgta agttacccac ttctggtgct agtttgtatg ttaatttgtt 60 140 tttatcctga tacatataat caggtagaat tgggaatttt attcagcctc cgcgtcgggg 120 141 aacatagaat aactotttat tqactoaaga otggaatttt cagaaaagtt taaattttta 180 142 ttttttctag aattttcaga aaagtttaat agatctgaga catttttaaa tcttttaatc 240 143 tttctagcta tttgtgaata tgcttttctt ccttttaaat aaatataata gctggtatgt 300 W--> 144 aaganagcta ttgatgcata tttttatttg gatatettat tgaactetta attggaataa 360 145 ttttcagt 148 <210> SEO ID NO: 7 149 <211> LENGTH: 583 150 <212> TYPE: DNA 151 <213> ORGANISM: Homo sapiens 153 <220> FEATURE: 154 <221> NAME/KEY: unsure 155 <222> LOCATION: (61)..(80) 156 <223> OTHER INFORMATION: a, c, g or t 158 <220> FEATURE: 159 <221> NAME/KEY: unsure 160 <222> LOCATION: (495) 161 <223> OTHER INFORMATION: a, c, g or t 163 <400> SEQUENCE: 7 164 cacaqcaqca tetqqaetqq tatetqaeca aaaactaqqt accatqqtet aqceacattt 60 W--> 165 nnnnnnnnn nnnnnnnnn atagtcataa gatttggttt tctcttactg aaatgcggta 120 166 taattgggat ttattcaaat cttcttttat atccatccca taaagttttg taatttttcc 180 167 ttattggcct aacaattcct gtaaagatta tttgtaagta gttaataatt tttgtaatca 240 168 tgatgagttt aacactttta ttttattaca tgtgctcata tgttgttgct gacactaaga 300 169 aacctatagg tcaaaagatg caaactaggg ccacatgagc acggcagcac ccagccagga 360 170 ctctgctgca gctgccgttt gtagatggag ctctctgtct ccaaagaagc acaggcctgt 420 171 tgttcttctg gttgtgctac agtaaaatga acctgggttt tctgaacatg tggttgaatg 480 W--> 172 tcagttgcta cctancgttt cacttgtgat attgatttta ttaatcttaa agttatgtga 540 173 atgttaacta tttctcatat ctatatctta ttcaaaatct ggg 583 176 <210> SEQ ID NO: 8

177 <211> LENGTH: 118

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Input Set : A:\Dex-249.app

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- 178 <212> TYPE: DNA
- 179 <213> ORGANISM: Homo sapiens
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- 183 tatgtttctt aattgtacca tgaattacaa aaacctattg gcaagatcag tcttattt 118
- 186 <210> SEQ ID NO: 9
- 187 <211> LENGTH: 502
- 188 <212> TYPE: DNA
- 189 <213> ORGANISM: Homo sapiens
- 191 <400> SEOUENCE: 9
- 192 tocacttttt aaaaagagtt agttatatgt agcatagtaa aaatatttct tttgaaatat 60
- 193 actattatat ccatttattt aagttctaat agactatact gcctaaaaaa atagagttga 120
- 194 tcttagtcaa atcacttaaa tcaggaacct ttaagattca gtttaatcag ttttgtatat 180
- 195 aaagtattag acttactttc accatttttg cccaaaaaca aacttctgta cttcattaaa 240
- 196 taacatcaac aaaagtaaaa agctaacatc aacctaggaa aaccattttc accgtaacag 300
- 197 ttatgtatat taactatatt atatactttt aaattagtaa aaaaaagggt aaaagacatg 360
- 198 aacaattatc aaaggaaaaa aatcctgaaa tattcaaatg tttaccttac taaaggaatt 420
- 199 cttaatctgc tgattatgtt tcttaattgt accatgaatt acaaaaacct attggcaaga 480
- 200 tcagtcttat ttaaaaaaaa aa
- 203 <210> SEQ ID NO: 10
- 204 <211> LENGTH: 390
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- 206 <213> ORGANISM: Homo sapiens
- 208 <220> FEATURE:
- 209 <221> NAME/KEY: unsure
- 210 <222> LOCATION: (83)..(211)
- 211 <223> OTHER INFORMATION: a, c, g or t
- 213 <220> FEATURE:
- 214 <221> NAME/KEY: unsure
- 215 <222> LOCATION: (214)
- 216 <223> OTHER INFORMATION: a, c, g or t
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- 219 <221> NAME/KEY: unsure
- 220 <222> LOCATION: (300)
- 221 <223> OTHER INFORMATION: a, c, g or t
- 223 <220> FEATURE:
- 224 <221> NAME/KEY: unsure
- 225 <222> LOCATION: (304)..(324)
- 226 <223> OTHER INFORMATION: a, c, g or t
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- 229 <221> NAME/KEY: unsure
- 230 <222> LOCATION: (368)
- 231 <223> OTHER INFORMATION: a, c, g or t
- 233 <220> FEATURE:
- 234 <221> NAME/KEY: unsure
- 235 <222> LOCATION: (381)
- 236 <223> OTHER INFORMATION: a, c, g or t
- 238 <400> SEQUENCE: 10
- 239 cacatatate cagtaacagg getgtgegtg taccetetaa atetacaaaa ataaaaagte 60

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W--> 242 nnnnnnnnn nnnnnnnnn nnnnnnnnn nacncgtatt cetttetgga gaetetaggg 240
W--> 243 qaatctcttt tcttgccctt tctggcttct agaacctgcc tacattcttt ggctagtggn 300
W--> 244 cccnnnnnn nnnnnnnnn nnnngtggct ggtcagtctt tctctgatgt tatctttctg 360
W--> 245 gttctgancc ttccatctcc ntcttcgcca
    248 <210> SEQ ID NO: 11
    249 <211> LENGTH: 266
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    251 <213> ORGANISM: Homo sapiens
    253 <400> SEQUENCE: 11
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    255 ggcaaaacct tgggagtttt gaggtccctg tggacttgcc tggcatgcct gggagatttg 120
    256 taggtatett etttetaeae ggattgtgtt tetgeeetea aggaaaatag tteaetttga 180
    257 ccactgtaaa tgatgtagta tttaaacaaa aggaaaagca cttcattgtc tgctctaaaa 240
    258 ctaaaatgtt aagaaagaga ggtggc
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    262 <211> LENGTH: 380
    263 <212> TYPE: DNA
    264 <213> ORGANISM: Homo sapiens
    266 <400> SEQUENCE: 12
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    268 gctggaaatt tggatacagt aaatggtata taaggcttct ttttgcttgt gcaccagaag 120
    269 tetttgtece ageeteeaga teageagtea gtacteeeet eteceaacet gtaggaagta 180
    270 cttgtgaaaa gttatctatc cctggtctga gtgggaggtt cttaacctca ttgatgtttt 240
    271 agtgtgactt gtctacattt gtgtgctccc ttcgtcatct gcagaggata tgagaaaaga 300
    272 aacaaatgaa caaaaagtgg agatagcgcc ttcctattca ttcttcattt ggtatggtta 360
    273 taqttaaqaq aqqtqaqcca
    276 <210> SEQ ID NO: 13
    277 <211> LENGTH: 871
    278 <212> TYPE: DNA
    279 <213> ORGANISM: Homo sapiens
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    283 ctcattgcag cctcggtctc ccaggctcaa gcaatccttc cacctcagcc tccagactag 120
    284 ctggaactac aggcgcatat caccaaqcct qqctaatttt ttaattaaqa caqqqtctac 180
    285 tatgtggccc aggetggtct tgaactcctg gactcaagca atteteccac tetggectec 240
    286 caaagtgata ggattacagg catgagccac cacgcccagc ctggctcacc tctcttaact 300
    287 ataaccatac caaatgaaga atgaatagga aggcgctatc tccacttttt gttcatttgt 360
    288 ttcttttctc atatcctctg cagatgacga agggagcaca caaatgtaga caagtcacac 420
    289 taaaacatca atgaggttaa gaacctccca ctcagaccag ggatagataa ctttcacagt 480
    290 acttctacag ttggagaggg agtactactc. tgatctggag gctggacaaa acttctgtgc 540
    291 acaagcaaaa aaagcctatt accattacct qtatccaaat tccagcagca acgatggcga 600
    292 agageeteat ageagatgtg geagtggeeg eeactteeea tgacaaaggg aggtaggeat 660
    293 gattcatete tagtgeatgg ggaeggttge ettttgeece aaateaacte aagaeceett 720
    294 tocaccatat gtotgtatgt aactotaaat goatototaa gaottaagaa taaaaagooa 780
    295 ctgctacctg gcaagtggat gaggcaaaag agaagacata ccccaaagaa ctatagcact 840
    296 ctgctccaaa ttacaqaact ttctaaacgt c
    299 <210> SEQ ID NO: 14
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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\Dex-249.app

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L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:240~M:341~W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:350~M:341~W:~(46)~"n" or "Xaa" used, for SEQ ID#:15
L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:352 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:387 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:521~M:341~W:~(46) "n" or "Xaa" used, for SEQ ID#:23
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:558 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:559 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:633~M:341~W:~(46)~"n"~or~"Xaa"~used, for SEQ~ID#:29
L:634 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:721 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:723 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:768 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:769 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:770~M:341~W:~(46) "n" or "Xaa" used, for SEQ ID#:38
L:871 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:922 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47
L:1081 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:1100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:1172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:1176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:1646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:1795 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:1849 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
L:1850 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
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L:1950 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92
L:1951 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92
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L:4236 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:177
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L:6168\ M:258\ W: Mandatory Feature missing, <223> not found for SEQ ID#:256
L:6171 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:256
L:6174 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:256
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L:6220 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:258
L:6237 \ M:258 \ W: Mandatory Feature missing, <223> not found for SEQ ID#:259
L:6240~M:258~W: Mandatory Feature missing, <223> not found for SEQ ID#:259
L:6243 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:259
L:6246 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:259
L:6768 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:264
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L:6780 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:264
L:7247 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:280
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